

TEAM 4

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**RAW SEQUENCE LISTING
PATENT APPLICATION US/08/657,749**

DATE: 07/26/96
TIME: 11:14:39

INPUT SET: S11781.raw

This Raw Listing contains the General Information Section and those Sequences containing ERRORS.

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/657,749DATE: 07/26/96
TIME: 11:14:42*INPUT SET: S11781.raw*

46
47 (vii) CURRENT APPLICATION DATA:
48
49 (A) APPLICATION NUMBER:
50
51 (B) FILING DATE:
52
53 (C) CLASSIFICATION:
54
55 (vii) PRIOR APPLICATION DATA:
56
57 (A) APPLICATION NUMBER: 08/265,047
58
59 (B) FILING DATE: 23-JUN-94
60
61 (C) CLASSIFICATION:
62
63 (vii) PRIOR APPLICATION DATA:
64
65 (A) APPLICATION NUMBER: 08/160,602
66
67 (B) FILING DATE: 30-NOV-93
68
69 (C) CLASSIFICATION:
70
71 (vii) PRIOR APPLICATION DATA:
72
73 (A) APPLICATION NUMBER: 08/066,299
74
75 (B) FILING DATE: 20-MAY-93
76
77 (vii) PRIOR APPLICATION DATA:
78
79 (A) APPLICATION NUMBER: PCT/US92/09863
80
81 (B) FILING DATE: 13-NOV-92
82
83 (vii) PRIOR APPLICATION DATA:
84
85 (A) APPLICATION NUMBER: 07/933,411
86
87 (B) FILING DATE: 21-AUG-92
88
89 (vii) PRIOR APPLICATION DATA:
90
91 (A) APPLICATION NUMBER: 07/796,256
92
93 (B) FILING DATE: 20-NOV-91
94
95 (viii) ATTORNEY/AGENT INFORMATION:
96
97 (A) NAME: Donna E. Scherer
98

**RAW SEQUENCE LISTING
PATENT APPLICATION US/08/657,749**

DATE: 07/26/96
TIME: 11:14:46

INPUT SET: S11781.raw

99 (B) REGISTRATION NUMBER: 34,719
100
101 (A) NAME: Carl J. Schwedler
102
103 (B) REGISTRATION NUMBER: 36,924
104
105 (C) REFERENCE/DOCKET NUMBER: CGNE 101-2 US
106
107 (ix) TELECOMMUNICATION INFORMATION:
108
109 (A) TELEPHONE: (916) 753-6313
110
111 (B) TELEFAX: (916) 753-1510
112
113
114
--> 115 (2) INFORMATION FOR SEQ ID NO: 1:
116
117 (i) SEQUENCE CHARACTERISTICS:
118
119 (A) LENGTH: 1786 base pairs
120
121 (B) TYPE: nucleic acid
122
123 (C) STRANDEDNESS: single
124
125 (D) TOPOLOGY: linear
126
127 (ii) MOLECULE TYPE: cDNA to mRNA
128
--> 129 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
130
131 AAATCCTCCA CTCATACT CCACCTCTCT CTCTCTCTCT CTCTCTGA AACAAATTG 60
132
133
134 GTAGCAAAC TAAAAAGAAA ATG GAG GAA ATG GGA AGC ATT TTA GAG TTT CTT 112
135 Met Glu Glu Met Gly Ser Ile Leu Glu Phe Leu
136 1 5 10
137
138
139 GAT AAC AAA GCC ATT TTG GTC ACT GGT GCT ACT GGC TCC TTA GCA AAA 160
140 Asp Asn Lys Ala Ile Leu Val Thr Gly Ala Thr Gly Ser Leu Ala Lys
141 15 20 25
142
143
144 ATT TTT GTG GAG AAG GTA CTG AGG AGT CAA CCG AAT GTG AAG AAA CTC 208
145 Ile Phe Val Glu Lys Val Leu Arg Ser Gln Pro Asn Val Lys Lys Leu
146 30 35 40
147
148
149 TAT CTT CTT TTG AGA GCA ACC GAT GAC GAG ACA GCT GCT CTA CGC TTG 256
150 Tyr Leu Leu Leu Arg Ala Thr Asp Asp Glu Thr Ala Ala Leu Arg Leu
151 45 50 55

**RAW SEQUENCE LISTING
PATENT APPLICATION US/08/657,749**DATE: 07/26/96
TIME: 11:14:49**INPUT SET: S11781.raw**

152
153
154 CAA AAT GAG GTT TTT GGA AAA GAG TTG TTC AAA GTT CTG AAA CAA AAT 304
155 Gln Asn Glu Val Phe Gly Lys Glu Leu Phe Lys Val Leu Lys Gln Asn
156 60 65 70 75
157
158
159 TTA GGT GCA AAT TTC TAT TCC TTT GTA TCA GAA AAA GTG ACT GTA GTA 352
160 Leu Gly Ala Asn Phe Tyr Ser Phe Val Ser Glu Lys Val Thr Val Val
161 80 85 90
162 CCC GGT GAT ATT ACT GGT GAA GAC TTG TGT CTC AAA GAC GTC AAT TTG 400
163 Pro Gly Asp Ile Thr Gly Glu Asp Leu Cys Leu Lys Asp Val Asn Leu
164 95 100 105
165
166
167 AAG GAA GAA ATG TGG AGG GAA ATC GAT GTT GTT GTC AAT CTA GCT GCT 448
168 Lys Glu Glu Met Trp Arg Glu Ile Asp Val Val Val Asn Leu Ala Ala
169 110 115 120
170
171
172 ACA ATC AAC TTC ATT GAA AGG TAC GAC GTG TCT CTG CTT ATC AAC ACA 496
173 Thr Ile Asn Phe Ile Glu Arg Tyr Asp Val Ser Leu Leu Ile Asn Thr
174 125 130 135
175
176
177 TAT GGA GCC AAG TAT GTT TTG GAC TTC GCG AAG AAG TGC AAC AAA TTA 544
178 Tyr Gly Ala Lys Tyr Val Leu Asp Phe Ala Lys Lys Cys Asn Lys Leu
179 140 145 150 155
180
181
182 AAG ATA TTT GTT CAT GTA TCT ACT GCT TAT GTA TCT GGA GAG AAA AAT 592
183 Lys Ile Phe Val His Val Ser Thr Ala Tyr Val Ser Gly Glu Lys Asn
184 160 165 170
185
186
187 GGG TTA ATA CTG GAG AAG CCT TAT TAT ATG GGC GAG TCA CTT AAT GGA 640
188 Gly Leu Ile Leu Glu Lys Pro Tyr Tyr Met Gly Glu Ser Leu Asn Gly
189 175 180 185
190
191
192 AGA TTA GGT CTG GAC ATT AAT GTA GAG AAG AAA CTT GTG GAG GCA AAA 688
193 Arg Leu Gly Leu Asp Ile Asn Val Glu Lys Lys Leu Val Glu Ala Lys
194 190 195 200
195
196
197 ATC AAT GAA CTT CAA GCA GCG GGG GCA ACG GAA AAG TCC ATT AAA TCG 736
198 Ile Asn Glu Leu Gln Ala Ala Gly Ala Thr Glu Lys Ser Ile Lys Ser
199 205 210 215
200
201
202 ACA ATG AAG GAC ATG GGC ATC GAG AGG GCA AGA CAC TGG GGA TGG CCA 784
203 Thr Met Lys Asp Met Gly Ile Glu Arg Ala Arg His Trp Gly Trp Pro
204 220 225 230 235

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/657,749DATE: 07/26/96
TIME: 11:14:53

INPUT SET: SII781.raw

205
206
207 AAT GTG TAT GTA TTC ACC AAG GCA TTA GGG GAG ATG CTT TTG ATG CAA 832
208 Asn Val Tyr Val Phe Thr Lys Ala Leu Gly Glu Met Leu Leu Met Gln
209 240 245 250
210
211
212 TAC AAA GGG GAC ATT CCG CTT ACT ATT ATT CGT CCC ACC ATC ATC ACC 880
213 Tyr Lys Gly Asp Ile Pro Leu Thr Ile Ile Arg Pro Thr Ile Ile Thr
214 255 260 265
215
216
217 AGC ACT TTT AAA GAG CCC TTT CCT GGT TGG GTT GAA GGT GTC AGG ACC 928
218 Ser Thr Phe Lys Glu Pro Phe Pro Gly Trp Val Glu Gly Val Arg Thr
219 270 275 280
220
221
222 ATC GAT AAT GTA CCT GTA TAT TAT GGT AAA GGG AGA TTG AGG TGT ATG 976
223 Ile Asp Asn Val Pro Val Tyr Tyr Gly Lys Gly Arg Leu Arg Cys Met
224 285 290 295
225
226
227
228 CTT TGC GGA CCC AGC ACA ATA ATT GAC CTG ATA CCG GCA GAT ATG GTC 1024
229 Leu Cys Gly Pro Ser Thr Ile Ile Asp Leu Ile Pro Ala Asp Met Val
230 300 305 310 315
231
232
233 GTG AAT GCA ACG ATA GTA GCC ATG GTG GCG CAC GCA AAC CAA AGA TAC 1072
234 Val Asn Ala Thr Ile Val Ala Met Val Ala His Ala Asn Gln Arg Tyr
235 320 325 330
236
237
238 GTA GAG CCG GTG ACA TAC CAT GTG GGA TCT TCA GCG GCG AAT CCA ATG 1120
239 Val Glu Pro Val Thr Tyr His Val Gly Ser Ser Ala Ala Asn Pro Met
240 335 340 345
241
242
243 AAA CTG AGT GCA TTA CCA GAG ATG GCA CAC CGT TAC TTC ACC AAG AAT 1168
244 Lys Leu Ser Ala Leu Pro Glu Met Ala His Arg Tyr Phe Thr Lys Asn
245 350 355 360
246
247
248 CCA TGG ATC AAC CCG GAT CGC AAC CCA GCA CAT GTG GGT CGG GCT ATG 1216
249 Pro Trp Ile Asn Pro Asp Arg Asn Pro Val His Val Gly Arg Ala Met
250 365 370 375
251
252
253 GTC TTC TCC TCC TTC ACC TTC CAC CTT TAT CTC ACC CTT AAT TTC 1264
254 Val Phe Ser Ser Phe Ser Thr Phe His Leu Tyr Leu Thr Leu Asn Phe
255 380 385 390 395
256
257

RAW SEQUENCE LISTING
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277
278 ATA GTT GAA GCT GAT ATG TTT TAC TTT GAT CCC AGG GCA ATT AAC TGG 1504
279 Ile Val Glu Ala Asp Met Phe Tyr Phe Asp Pro Arg Ala Ile Asn Trp
280 460 465 470 475
281
282
283
284 GAA GAT TAC TTC TTG AAA ACT CAT TTC CCA GGN GTC GTA GAG CAC GTT 1552
285 Glu Asp Tyr Phe Leu Lys Thr His Phe Pro Gly Val Val Glu His Val
286 480 485 490
287
288
289 CTT AAC TAAAAGTTAC GGTACGAAAA TGAGAAGATT GGAATGCATG CACCGAAAGN 1608
290 Leu Asn
291
292
293 NCAACATAAA AGACGTGGTT AAAGTCATGG TCAAAAAAGA AATAAAATGC AGTTAGGTTT 1668
294
295
296 GTGTTGCAGT TTTGATTCCCT TGTATTGTTA CTTGTACTTT TGATCTTTTT CTTTTTTAAT 1728
297
298
299 GAAATTTCTC TCTTGTTTT GTGAAAAAAA AAAAAAAA GAGCTCCTGC AGAAGCTT 1786
300
301

986 (2) INFORMATION FOR SEQ ID NO: 15:

987
988 (i) SEQUENCE CHARACTERISTICS:
989
990 (A) LENGTH: 17 base pairs
991
992 (B) TYPE: nucleic acid
993
994 (C) STRANDEDNESS: single
995
996 (D) TOPOLOGY: linear
997
998 (ii) MOLECULE TYPE: other
999

--> 1000 (A) synthetic oligonucleotide

1001
1002 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15 :

1003
1004 AAYATHACNA CNYTNNGG

17

1009 (2) INFORMATION FOR SEQ ID NO: 16:

1010
1011 (i) SEQUENCE CHARACTERISTICS:

Desr. pt. or

RAW SEQUENCE LISTING
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1012
1013 (A) LENGTH: 17 base pairs
1014
1015 (B) TYPE: nucleic acid
1016
1017 (C) STRANDEDNESS:single
1018
1019 (D) TOPOLOGY: linear
1020
1021 (ii) MOLECULE TYPE: other
1022
--> 1023 (A) synthetic oligonucleotide
1024
1025 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16 :
1026
1027
1028 SWRTTRCAYT TRAANCC 17
1029
1030
1031

2114 (2) INFORMATION FOR SEQ ID NO: 25:
2115
2116 (i) SEQUENCE CHARACTERISTICS:
2117
2118 (A) LENGTH: 1664 base pairs
2119
2120 (B) TYPE: nucleic acid
2121
2122 (C) STRANDEDNESS: single
2123
2124 (D) TOPOLOGY: linear
2125
2126 (ii) MOLECULE TYPE: cDNA to mRNA
2127
2128
2129
2130
2131 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:
2132
2133
2134 CA ATG ACG TCT GTG AAC GTA AAA CTC CTT TAC CAT TAC GTC ATA ACC 47
2135 Met Thr Ser Val Asn Val Lys Leu Leu Tyr His Tyr Val Ile Thr
2136 1 5 10 15
2137
2138
2139 AAC TTT TTC AAC CTC TGT TTC TTC CCA CTG ACG GGG ATC CTC GCC GGA 95
2140 Asn Phe Phe Asn Leu Cys Phe Phe Pro Leu Thr Gly Ile Leu Ala Gly
2141 20 25 30
2142
2143
2144 AAA GGC TCT CGT CTT ACC ACA AAC GAT CTC CAC CAC TTC TAT TCA TAT 143
2145 Lys Gly Ser Arg Leu Thr Thr Asn Asp Leu His His Phe Tyr Ser Tyr

**RAW SEQUENCE LISTING
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2146	35	40	45	
2147				
2148				
2149	CTC CAA CAC AAN CTT ATA ACC TTA ACC CTA CTC TTT GGC TTC ACC GTT			191
2150	Leu Gln His Xxx Leu Ile Thr Leu Thr Leu Leu Phe Gly Phe Thr Val			
2151	50	55	60	
2152				
2153				
2154	TTT GGT TCG GTT CTC TAC TTC GTA ANC CGA CCC AAA CCG GTT TAC CTC			239
2155	Phe Gly Ser Val Leu Tyr Phe Val Xxx Arg Pro Lys Pro Val Tyr Leu			
2156	65	70	75	
2157				
2158				
2159	GTT GAC TAC TCC TGC TAC CTT CCA CCA CAA CAT CTT AGC GCT GGT ATC			287
2160	Val Asp Tyr Ser Cys Tyr Leu Pro Pro Gln His Leu Ser Ala Gly Ile			
2161	80	85	90	95
2162				
2163				
2164	TCT AAG ACC ATG GAA ATC TTT TAT CAA ATA AGA AAA TCT GAT CCT TTA			335
2165	Ser Lys Thr Met Glu Ile Phe Tyr Gln Ile Arg Lys Ser Asp Pro Leu			
2166	100	105	110	
2167				
2168				
2169	CGA AAC GTG GCA TTA GAT GAT TCG TCT TCT GAT TTC TTG AGA AAG			383
2170	Arg Asn Val Ala Leu Asp Asp Ser Ser Ser Leu Asp Phe Leu Arg Lys			
2171	115	120	125	
2172				
2173				
2174	ATT CAA GAG CGT TCA GGT CTA GGC GAT GAA ACC TAC GGC CCC GAG GGA			431
2175	Ile Gln Glu Arg Ser Gly Leu Gly Asp Glu Thr Tyr Gly Pro Glu Gly			
2176	130	135	140	
2177				
2178				
2179	CTG TTT GAG ATT CCT CCG AGG AAG AAT TTA GCG TCG GCG CGT GAA GAG			479
2180	Leu Phe Glu Ile Pro Pro Arg Lys Asn Leu Ala Ser Ala Arg Glu Glu			
2181	145	150	155	
2182				
2183				
2184				
2185				
2186	ACG GAG CAA GTA ATC AAC GGT GCG CTA AAA AAT CTA TTC GAG AAC AAC			527
2187	Thr Glu Gln Val Ile Asn Gly Ala Leu Lys Asn Leu Phe Glu Asn Asn			
2188	160	165	170	175
2189				
2190				
2191	AAA GTT AAC CCT AAA GAG ATT GGT ATA CTT GTG GTG AAC TCA AGC ATG			575
2192	Lys Val Asn Pro Lys Glu Ile Gly Ile Leu Val Val Asn Ser Ser Met			
2193	180	185	190	
2194				
2195				
2196	TTT AAT CCG ACT CCT TCG TTA TCC GCG ATG GTA GTT AAT ACT TCC AAG			623
2197	Phe Asn Pro Thr Pro Ser Leu Ser Ala Met Val Val Asn Thr Ser Lys			
2198	195	200	205	

**RAW SEQUENCE LISTING
PATENT APPLICATION US/08/657,749**

 DATE: 07/26/96
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2199			
2200			
2201	CTC CGA AGC AAC ATC AAA AGC TTT AAT CTT GGA GGA ATG GGT TGC AGT		671
2202	Leu Arg Ser Asn Ile Lys Ser Phe Asn Leu Gly Gly Met Gly Cys Ser		
2203	210	215	220
2204			
2205			
2206	GCT GGT GTT ATC GCC ATT GAT CTA GCT AAA GAC TTG TTG CAT GTT CAT		719
2207	Ala Gly Val Ile Ala Ile Asp Leu Ala Lys Asp Leu Leu His Val His		
2208	225	230	235
2209			
2210			
2211	AAA AAC ACA TAT GCT CTT GTG GTG AGC ACA GAG AAC ATC ACT CAA AAC		767
2212	Lys Asn Thr Tyr Ala Leu Val Val Ser Thr Glu Asn Ile Thr Gln Asn		
2213	240	245	250
2214			
2215			
2216	ATT TAT ACC GGT GAT AAC AGA TCC ATG ATG GTT TCG AAT TGC TTG TTC		815
2217	Ile Tyr Thr Gly Asp Asn Arg Ser Met Met Val Ser Asn Cys Leu Phe		
2218	260	265	270
2219			
2220			
2221	CGT GTC GGT GGG GCA GCG ATT CTG CTC TCC AAC AAG CCG GGG GAT CGA		863
2222	Arg Val Gly Gly Ala Ala Ile Leu Leu Ser Asn Lys Pro Gly Asp Arg		
2223	275	280	285
2224			
2225			
2226	AGA CGG TCC AAG TAC AAG CTA GCT CAC ACG GTT CGA ACG CAT ACC GGA		911
2227	Arg Arg Ser Lys Tyr Lys Leu Ala His Thr Val Arg Thr His Thr Gly		
2228	290	295	300
2229			
2230			
--> 2231	GCT GAC GAC AAG TCT TTT GGA TGT GTG CGG CAA GAA GAA GAT GAT AGC		949
2232	Ala Asp Asp Lys Ser Phe Gly Cys Val Arg Gln Glu Glu Asp Asp Ser		
2233	305	310	315
2234			
2235			
--> 2236	GGT AAA ACC GGA GTT AGT TTG TCA AAA GAC ATA ACC GTT GTT GCC GGG		1007
2237	Gly Lys Thr Gly Val Ser Leu Ser Lys Asp Ile Thr Val Val Ala Gly		
2238	320	325	330
2239			
2240			
2241			
2242	ATA ACG GTT CAG AAA AAC ATA ACA ACA TTG GGT CCG TTG GTT CTT CCT		1055
2243	Ile Thr Val Gln Lys Asn Ile Thr Thr Leu Gly Pro Leu Val Leu Pro		
2244	340	345	350
2245			
2246			
2247	CTG AGC GAA AAA ATC CTT TTT GTC GTT ACA TTC GTA GCC AAG AAA CTA		1103
2248	Leu Ser Glu Lys Ile Leu Phe Val Val Thr Phe Val Ala Lys Lys Leu		
2249	355	360	365
2250			
2251			

edit ↴

949 ↵ OK

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/657,749DATE: 07/26/96
TIME: 11:15:23

INPUT SET: S11781.raw

2252 TTA AAA GAT AAG ATC AAA CAC TAT TAC GTG CCG GAT TTC AAA CTT GCA 1151
2253 Leu Lys Asp Lys Ile Lys His Tyr Tyr Val Pro Asp Phe Lys Leu Ala
2254 370 375 380
2255
2256
2257 GTA GAT CAT TTC TGT ATT CAT GCG GGA GGT AGA GCC GTG ATA GAT GTG 1199
2258 Val Asp His Phe Cys Ile His Ala Gly Gly Arg Ala Val Ile Asp Val
2259 385 390 395
2260
2261
2262 TTA GAG AAG AAC TTA GGG CTA TCG CCG ATA GAT GTG GAG GCA TCA AGA 1247
2263 Leu Glu Lys Asn Leu Gly Leu Ser Pro Ile Asp Val Glu Ala Ser Arg
2264 400 405 410 415
2265
2266
2267 TCA ACA TTA CAT AGA TTT GGG AAT ACA TCG TCT AGT TCA ATT TGG TAT 1295
2268 Ser Thr Leu His Arg Phe Gly Asn Thr Ser Ser Ser Ser Ile Trp Tyr
2269 420 425 430
2270
2271
2272 GAA TTA GCA TAC ATA GAG CCA AAA GGA AGG ATG AAG AAA GGT AAT AAA 1343
2273 Glu Leu Ala Tyr Ile Glu Pro Lys Gly Arg Met Lys Lys Gly Asn Lys
2274 435 440 445
2275
2276
2277 GCT TGC CAA ATA GCT GGT GGG TCA GGT TTT AAG TGT AAT AGT GCG GTT 1391
2278 Ala Cys Gln Ile Ala Gly Gly Ser Gly Phe Lys Cys Asn Ser Ala Val
2279 450 455 460
2280
2281
2282 TGG GTC GCT TTA CGC AAT GTC GAG GCT TCA GCT AAT AGT CCT TGG GAA 1439
2283 Trp Val Ala Leu Arg Asn Val Glu Ala Ser Ala Asn Ser Pro Trp Glu
2284 465 470 475
2285
2286
2287 CAT TGC ATT CAC AAA TAT CCG GTT CAA ATG TAT TCT GGT TCA TCA AAG 1487
2288 His Cys Ile His Lys Tyr Pro Val Gln Met Tyr Ser Gly Ser Ser Lys
2289 480 485 490 495
2290
2291
2292 TCA GAG ACT CCT GTC CAA AAC GGT CGG TCC TAATTTATGT ATCTCAAATG 1537
2293 Ser Glu Thr Pro Val Gln Asn Gly Arg Ser
2294 500
2295
2296
2297
2298 ATGTTGTCCA CTTTCTCTTT TTTTTTTCT TTTTTTAGTT ATAATTAAAT GGTTACGATG 1597
2299
2300
2301 TTTTGTCTAG GTCGTTATAA ATAAAGAATA CATGGGTGTT ACTAGTATAA AAAAAAAA 1657
2302
2303
2304 AAAAAAA 1664

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/657,749DATE: 07/26/96
TIME: 11:15:27

INPUT SET: S11781.raw

2305
2306

2627 (2) INFORMATION FOR SEQ ID NO: 30:

2628

2629 (i) SEQUENCE CHARACTERISTICS:

2630

2631 (A) LENGTH: 42 base pairs

2632

2633 (B) TYPE: nucleic acid

2634

2635 (C) STRANDEDNESS:single

2636

2637 (D) TOPOLOGY: linear

2638

2639 (ii) MOLECULE TYPE: other

2640

--> 2641 (A) synthetic oligonucleotide

2642

2643

2644 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:

2645

2646 CAUCAUCAUC AUGTCGACAA AATGACGTCC ATTAACGTAA AG

42

2647

2648

2649

2650 (2) INFORMATION FOR SEQ ID NO: 31:

2651

2652 (i) SEQUENCE CHARACTERISTICS:

2653

2654 (A) LENGTH: 50 base pairs

2655

2656 (B) TYPE: nucleic acid

2657

2658 (C) STRANDEDNESS:single

2659

2660 (D) TOPOLOGY: linear

2661

2662 (ii) MOLECULE TYPE: other

2663

--> 2664 (A) synthetic oligonucleotide

2665

2666

2667 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 31 :

2668

2669 CUACUACUAC UAGTCGACGG ATCCTATTTG GAAGCTTGA CATTGTTTAG

50

2670

2671

2672

2673

2696 (2) INFORMATION FOR SEQ ID NO: 33:

Same error

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/657,749DATE: 07/26/96
TIME: 11:15:30

INPUT SET: S11781.raw

2697
2698 (i) SEQUENCE CHARACTERISTICS:
2699
2700 (A) LENGTH: 41 base pairs
2701
2702 (B) TYPE: nucleic acid
2703
2704 (C) STRANDEDNESS:single
2705
2706 (D) TOPOLOGY: linear
2707
2708 (ii) MOLECULE TYPE: other
2709
--> 2710 (A) synthetic oligonucleotide
2711
2712 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 33 :
2713
2714 CAUCAUCAUC AUGAATTCAA GCTTAARYTN BKNTAYCAYT A 41
2715
2716
2717

2738 (2) INFORMATION FOR SEQ ID NO: 35:
2739
2740 (i) SEQUENCE CHARACTERISTICS:
2741
2742 (A) LENGTH: 41 base pairs
2743
2744 (B) TYPE: nucleic acid
2745
2746 (C) STRANDEDNESS:single
2747
2748 (D) TOPOLOGY: linear
2749
2750 (iii) MOLECULE TYPE: other
2751
--> 2752 (A) synthetic oligonucleotide
2753
2754 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 35 :
2755
2756 CAUCAUCAUC AUGAATTCAA GCTTAAYYTN GGNGGNATGG G 41
2757
2758
2759

2760 (2) INFORMATION FOR SEQ ID NO: 36:
2761
2762 (i) SEQUENCE CHARACTERISTICS:
2763
2764 (A) LENGTH: 40 base pairs
2765
2766 (B) TYPE: nucleic acid
2767

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2768 (C) STRANDEDNESS:single
2769 (D) TOPOLOGY: linear
2770 (ii) MOLECULE TYPE: other
2771
2772
2773
--> 2774 (A) synthetic oligonucleotide
2775
2776 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 36 :
2777
2778 CUACUACUAC UAGGATCCGT CGACCCATNC CNCCNARRTT 40
2779
2780



2800 (2) INFORMATION FOR SEQ ID NO: 38:
2801
2802 (i) SEQUENCE CHARACTERISTICS:
2803
2804 (A) LENGTH: 41 base pairs
2805
2806 (B) TYPE: nucleic acid
2807
2808 (C) STRANDEDNESS:single
2809
2810 (D) TOPOLOGY: linear
2811
2812 (ii) MOLECULE TYPE: other
2813
--> 2814 (A) synthetic oligonucleotide
2815
2816 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 38 :
2817
2818 CUACUACUAC UAGGATCCGT CGACSWRTTR CAYTTRAANC C 41
2819
2820
2821



2822 (2) INFORMATION FOR SEQ ID NO: 39:
2823
2824 (i) SEQUENCE CHARACTERISTICS:
2825
2826 (A) LENGTH: 29 base pairs
2827
2828 (B) TYPE: nucleic acid
2829
2830 (C) STRANDEDNESS:single
2831
2832 (D) TOPOLOGY: linear
2833
2834 (ii) MOLECULE TYPE: other
2835
--> 2836 (A) synthetic oligonucleotide
2837



08/657,749

Notice of Availability

Applicant Aid for Biotechnology Computer Readable Form (CRF) Sequence Listings Submissions

The Patent and Trademark Office (PTO) has developed a computer program, called Checker, that will aid applicants in identifying and correcting errors prior to making submissions for compliance with the Requirements for Patent Applications Containing Nucleotide Sequence and/or Amino Acid Sequence Disclosures (sequence rules: 37 CFR 1.821 through 1.825). (Final rules were published in the *Federal Register* (55 FR 18230) on May 1, 1990, and in the *PTO Official Gazette* (1114 Off.Gaz.PatOffice 29) on May 15, 1990.)

Checker is a DOS-based software program that is intended to assist users in determining whether errors may be present in the sequence listings, and is not intended to guarantee that the submission is error-free.

The most current version of the software will be available via computer downloading (details below). Copies on diskette are also available. Updated software versions will not be automatically mailed out; any updates will be announced in the *PTO Official Gazette*.

The software can be accessed/requested in the following locations:

- 1) Dial-up access to the Patent and Trademark Office Bulletin Board System.
Phone number: 703-305-8950
Cost: Free-of-charge
- 2) Dial-up access through the Internet. FTP site: ftp.uspto.gov
Login as "anonymous". Software is in directory /pub/checker
Cost: Free-of-charge
- 3) For diskette copies, telephone requests to 703-306-2600.
Cost: \$25.00

For Further Information Contact: Meredith Beckhardt at 703-308-4212.

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**RAW SEQUENCE LISTING
PATENT APPLICATION US/08/657,749**

DATE: 07/26/96
TIME: 11:15:37

INPUT SET: S11781.raw

2838 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 39 :
2839
2840 CUACUACUAC UASWRTTRCA YTTRAANCC
2841

29